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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/965,703

DATE: 10/18/2001
TIME: 16:19:41

Input Set : A:\A01020B.ST25.txt
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3 <110> APPLICANT: Rohm and Haas Company
4 Palli, Subba Reddy
5 Kapitskaya, Marianna Zinovjevna
6 Cress, Dean Ervin
8 <120> TITLE OF INVENTION: Novel Ecdysone Receptor-Based Inducible Gene Expression

System

10 <130> FILE REFERENCE: A01020B
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/965,703

13 <141> CURRENT FILING DATE: 2001-09-26
15 <150> PRIOR APPLICATION NUMBER: 60/191,355
16 <151> PRIOR FILING DATE: 2000-03-22
18 <150> PRIOR APPLICATION NUMBER: 60/269,799
19 <151> PRIOR FILING DATE: 2001-02-20
21 <150> PRIOR APPLICATION NUMBER: PCT/US01/09050
22 <151> PRIOR FILING DATE: 2001-03-21
24 <160> NUMBER OF SEQ ID NOS: 75
26 <170> SOFTWARE: PatentIn version 3.1
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 1288
30 <212> TYPE: DNA

31 <213> ORGANISM: Choristoneura fumiferana
33 <220> FEATURE:
34 <221> NAME/KEY: misc_feature
35 <223> OTHER INFORMATION: Novel Sequence
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45 cgacggaaat	gccaggagtg	ccgcctgaag	aagtgttag	ctgttaggcatt	gaggcctgag	240
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53 gacaagctgt	tggagacaaa	ccggcagaaa	aacatcccc	agttgacagc	caaccagcag	480
55 ttcccttatcg	ccaggctcat	ctggtaccag	gacgggtacg	agcagccctc	tgatgaagat	540
57 ttgaagagga	ttacgcagac	gtggcagcaa	gcccgcgtatg	aaaacgaaga	gtctgacact	600
59 ccctccgc	agatcacaga	gatgactatac	ctcacggtcc	aacttatcgt	ggagttcg	660
61 aagggattgc	cagggttcgc	caagatctcg	cagcctgatc	aaattacgt	gtttaaggct	720
63 tgctcaagtg	aggtaatgt	gctccgagtc	gcgcgacgt	acgatgcggc	ctcagacagt	780
65 gttctgttgc	cgaacaacca	agcgtacact	cgcgcacaact	acccgcaaggc	tggcatggcc	840
67 tacgtcatcg	aggatctact	gcacttctgc	cggtgcgtatgt	actctatggc	gttgacaaac	900
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71 ccgcaactgg	tggaaagaaat	ccagcggtac	tacctgaata	cgctccgc	ctatatcc	1020
73 aaccagctga	gcgggtcggc	gcgttcgtcc	gtcatatacg	gcaagatcct	ctcaatcc	1080
75 tctgagctac	gcacgctcg	catgcacaaac	tccaaatgt	gatctccct	caagctcaag	1140
77 aacagaaagc	tgccgcctt	cctcgaggag	atctggatg	tggcggacat	gtcgacacc	1200
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84 <210> SEQ ID NO: 2

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 101 ctgcctgtca gcacgacgac ggtggacgac cacatgccgc ccattatgca gtgtgaacct 240
 103 ccacccctg aagcagcaag gattcacgaa gtggcccaa gtttctctc cgacaagctg 300
 105 ttggagacaa accggcagaa aaacatcccc cagttgacag ccaaccagca gttccttata 360
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 119 gaggatctac tgcaactctg cccgtgcatt tactctatgg cggtggacaa catccattac 780
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 127 cgcacgctcg gcatgcggaaa ctccaacatg tgcatctccc tcaagctcaa gaacagaaaag 1020
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 141 <223> OTHER INFORMATION: Novel Sequence
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 149 atgcagtgtt aacccatccacc tcctgaagca gcaaggattt acgaagggtt cccaaagggtt 180
 151 ctctccgaca agctgttggaa gacaaaccgg cagaaaaaca tcccccaattt gacagccaa 240
 153 cagcgttcc ttatcgccag gtcacatcttgg taccaggacg ggtacgagca gccttctgtat 300
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 157 gacactccctt tccggccagat cacagagatg actatcctca cggtccaactt tatctggag 420
 159 ttcgcgaagg gattgccagg gttcgccaaat atctcgacg ctgatcaat tacctgtttt 480
 161 aaggcttgcgtaa caagtggatggtt aatgtatgcgc cggatcgcc gacgatacga tgccgcctca 540
 163 gacagtgttc tggtcgccaa caaccaaggc tacactcgcc acaactaccg caaggcttgc 600
 165 atggcctacg tcatacgagga tctactgcac ttctggcggtt gcatgtactt tatggcggtt 660
 167 gacaacatcc attacgcgtt gtcacggctt gtcgtcatctt ttctgtaccg gccagggtt 720
 169 gaggcagccgc aactgggttggaa agaaaatccag cggtaactacc tgaatacgctt cccgatctat 780
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175	ctcaagaaca	gaaagctgcc	gccttcctc	gaggagatct	ggatgtggc	ggacatgtcg	960
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197	actatcctca	cggtccaaact	tatcggtggag	ttcgcgaagg	gattgccagg	gttcgccaag	180
199	atctcgcagc	ctgatcaaataat	tacgctgttt	aaggcttgc	caagtggat	aatgatgctc	240
201	cgagtcgcgc	gacgatacga	tgcggcctca	gacagtgttc	tgttcgcgaa	caaccaagcg	300
203	tacactcgcg	acaactaccg	caaggctggc	atggctacg	tcatcgagga	tctactgcac	360
205	ttctgccgtt	gcatgtactc	tatggcgtt	gacaacatcc	attacgcgt	gctcacggct	420
207	gtcgtcatct	tttctgaccg	gccagggtt	gagcagccgc	aactggtgg	agaaatccag	480
209	cggtactacc	tgaatacgtct	ccgcatactat	atcctgaacc	agctgaggcg	gtcggcgcgt	540
211	tcgtccgtca	tatacggcaa	gatcctctca	atcctctgt	agctacgcac	gctcggcatg	600
213	caaaaactcca	acatgtgcat	ctccctcaag	ctcaagaaca	gaaagctgcc	gccttcctc	660
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226	<221>	NAME/KEY:	misc_feature				
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237	ctctccgaca	agctgttgaa	gacaaaccgg	cagaaaaaca	tcccccaatt	gacagccaaac	240
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247	aaggcttgc	caagtggat	aatgatgctc	cgatcgccgc	gacgatacga	tgccgcctca	540
249	gacagtgttc	tgttcgcgaa	caaccaagcg	tacactcgcg	acaactaccg	caaggctggc	600
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257	atccctgaacc	agctgagcg	gtcggcgcgt	tcgtccgtca	tatacggcaa	gatcctctca	840
259	atccctctgt	agctacgcac	gctcggcatg	caaaaactcca	acatgtgcat	ctccctcaag	900
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427	ttggcctctg	gtggcgccca	agactttgtt	aagaaggaga	ttcttgaccc	tatgacatgc	180
429	gagccgcggcc	agcatgccac	tattccgcata	ctacccgtat	aaatattggc	caagtgtcaa	240
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433	taccaggatg	gctatgagca	gccatctgaa	gaggatctca	ggcgatataat	gagtcaaccc	360
435	gatgagaacg	agagccaaac	ggacgtcagc	tttccgcata	taaccggat	aaccataactc	420
437	acggtccagt	tgattgttgc	gtttgttataa	ggtctaccag	cgtttacaaa	gatacccccag	480
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443	gattcttaca	aaatggccgg	aatggctgtat	aacattgaag	acctgtcga	tttccgcgc	660
445	caaattttct	cgatgaaggt	ggacaacgtc	gaatacgcgc	tttccactgc	cattgtgatc	720
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L:12 M:270 C: Current Application Number differs, Replaced Current Application Number